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<p>Background:</p> <p>In recent years, research on dimensional models of psychopathology has substantially increased. Some of the most promising models have included a dimension of general psychopathology that is often called the p factor. However, the research on the practical uses of such models is still scarce, as is the evaluation of models beyond their goodness-of-fit indexes. This study aims to contribute to filling this gap by examining the underlying structure of preadolescent psychopathology and answering an important practical question: whether such models of preadolescent psychopathology can be used in predicting substance use in adolescence, which is a critical period in the development of substance use disorders. In addition, the model containing the p factor is evaluated with methods that have been seldom used but that can provide a more nuanced picture of psychopathology.</p> <p>Methods:</p> <p>Using the data from the UK Household longitudinal study, the underlying structure of psychopathology was first modelled with three confirmatory factor analyses ($n = 3437$). The models in question were a two-factor model consisting of internalizing and externalizing; a one-factor model consisting of the p factor; and a bifactor model consisting of internalizing, externalizing and the p factor. The bifactor model was also evaluated with several bifactor model indices. Then the models that exhibited at least acceptable statistical fit were used to predict the use of several substances in adolescence ($n = 1610$).</p> <p>Results and conclusions:</p> <p>The two-factor model and the bifactor provided an acceptable fit to the data. Both the externalizing factor in the two-factor model and the p factor in the bifactor model were also able to predict the use of all substances. However, the bifactor model had issues with interpretation and contrary to theorizing, it could be characterized as primarily unidimensional. Future research is encouraged to compare the practical utilities of different models of psychopathology and to evaluate bifactor models in more detail.</p>			
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<p>Tausta:</p> <p>Viime vuosina psykopatologian dimensionaalisia malleja koskeva tutkimus on lisääntynyt runsaasti. Jotkut kaikkein lupaavimmista malleista ovat sisältäneet yleisen psykopatologian ulottuvuuden, jota kutsutaan usein p-faktoriksi. Tutkimus tällaisten mallien käytännön hyödyistä on kuitenkin vielä vähäistä, kuten on myös mallien arvioiminen muuten kuin sopivuusindekseillä. Tämä tutkimus tähtää täyttämään tutkimuskirjallisuuden aukkoa tarkastelemalla varhaisnuorten latenttia psykopatologiaa ja vastaamalla tärkeään käytännön kysymykseen: voidaanko varhaisnuorten psykopatologian malleja hyödyntää päihteidenkäytön ennustamisessa nuoruusiässä, joka on kriittinen vaihe päihdeongelmien kehityksessä. Lisäksi mallia, joka sisältää p-faktorin arvioidaan menetelmillä, joita ei olla käytetty paljoa, mutta jotka voivat auttaa muodostamaan yksityiskohtaisempaa kuvaa psykopatologiasta.</p> <p>Menetelmät:</p> <p>Tutkimuksessa hyödynnettiin aineistoa UK household longitudinal study -hankkeesta ja mallinnettiin ensin psykopatologian rakennetta kolmella konfirmatorisella faktorianalyysillä ($n = 3437$). Kyseiset mallit olivat kaksifaktorinen malli, joka koostui internalisoinnista ja eksternalisoinnista; yksifaktorinen malli, joka koostui p-faktorista; ja bifaktorimalli, joka koostui internalisoinnista, eksternalisoinnista ja p-faktorista. Bifaktorimallia myös arvioitiin useilla bifaktorimallien arviointiin tarkoitetuilla menetelmillä. Sitten malleilla, jotka olivat sopivuusindekseiltään vähintään hyväksyttäviä, ennustettiin erilaisten päihteiden käyttöä nuoruusiässä ($n = 1610$).</p> <p>Tulokset ja johtopäätökset:</p> <p>Kaksifaktorimalli ja bifaktorimalli sopivat aineistoon hyväksyttävällä tavalla. Sekä eksternalisointifaktori kaksifaktorimallissa että p-faktori bifaktorimallissa myös ennustivat kaikkien tutkittujen päihteiden käyttöä. Bifaktorimallissa oli kuitenkin tulkinnallisia ongelmia ja teoretisoinnin vastaisesti sitä voisi kuvailla pääasiassa yksiulotteiseksi. Tulevaa tutkimusta kannustetaan vertailemaan psykopatologian eri mallien käytännön hyödynnettävyyksiä ja arvioimaan bifaktorimalleja aiempaa yksityiskohtaisemmin.</p>			
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Preadolescent general psychopathology and its use in prediction of substance use in adolescence

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Contents

1. Introduction.....	1
1.1. Background Assumptions of Categorical Classifications of Mental Disorders.....	1
1.2. The Proposed Latent Structures of Psychopathology	4
1.2.1. The general factor of psychopathology.	5
1.3. Substance Use	8
1.4. Research Questions	9
2. Methods	10
2.1. Participants.....	10
2.2. Measures	10
2.3. Procedure	11
2.4. Statistical Analyses	11
2.4.1. Confirmatory factor analyses, model fit and properties of the models.	12
2.4.2. Structural equation models.....	13
3. Results.....	15
3.1. Confirmatory Factor Analyses and Model Statistics	15
3.2. Structural Equation Models and Prediction of Substance Use	18
4. Discussion.....	20
4.1. Limitations	23
4.2. Conclusion	24
5. References.....	26

1. Introduction

In recent years, research on latent factors underlying all psychopathology has increased substantially along with discontentment towards categorical classifications. As a result, several competing dimensional models of latent structure of psychopathology have emerged. One of the most promising developments has been the surfacing of the so-called general psychopathology factor (p factor) – a general factor analogous to the general intelligence factor (g).

So far research concerning the p factor has examined in depth which factor models are viable in terms of statistical fit. It has been found that a model including a p factor can be applied to adolescents (e.g. Carragher et al., 2016; Caspi et al., 2014; Gomez, Stavropoulos, Vance, & Griffiths, 2019; Lahey et al., 2012) and possibly to preadolescents or children as well (Gomez et al., 2019; Lahey et al., 2015; Martel et al., 2016; Olino et al., 2018). However, fit indexes alone cannot determine whether a model is viable or useful. When trying to figure out the most appropriate way to model psychopathology, research should also examine other important model properties, such as the replicability of factors, and the practical uses of the models.

One area where models of underlying psychopathology could prove practical lies in predicting negative outcomes before they even occur. For example, adolescence is a critical period in the development of substance use disorders. Early substance use is a significant risk factor for later substance use disorders (Jordan & Andersen, 2017), which are in many ways burdensome for the individual and share a high comorbidity with other mental disorders (Kessler, Chiu, Demler, & Walters, 2005). Thus, knowledge of risk factors could be used for early prevention and intervention.

The aim of this study is to examine the viability and practicality of models underlying adolescent psychopathology. The statistical fit of several factor models of psychopathology is investigated. Then the models with at least acceptable fit are used to predict the use of various substances approximately three years later in adolescence.

1.1. Background Assumptions of Categorical Classifications of Mental Disorders

Traditionally mental disorders have been classified as categorical, discrete and assumedly distinct entities, which can be diagnosed on the basis of symptoms. This practice is in line with the standard nosological style of the rest of medicine, in which disorders are conceptualized as dichotomous entities, which either affect a person or do not.

The standard categorical approach remains highly influential to this day. The classifications in the current diagnostic manuals, The Diagnostic and Statistical Manual of Mental Disorders (5th ed.; DSM-5; American Psychiatric Association, 2013) and International Classification of Diseases (10th ed.; ICD-10; World Health Organization, 1992), provide clinicians, researchers and the general public alike with crucial tools for practice and thinking. For example, for clinicians and researchers discrete categories serve as means to classify complex phenomena and convey information about them in a systematic and generally agreed way. In folk psychology diagnostic classes have made their way into everyday speech in explaining aberrant behaviour. In some cases, being classified as either having or not having a mental disorder can even have judicial consequences.

However, despite their wide usage, categorical classifications of mental disorders have been a subject to a wide array of criticism (see Carragher, Krueger, Eaton, & Slade, 2015 for review). In short, most issues are related to the validity of such classifications. The commonly set objectives for classifications of mental disorders is that they should accurately describe how the human psychopathology functions or at least serve as instrumentally effective tools in practice (Zachar & Kendler, 2010). However, several flaws in the core assumptions of DSM and ICD might make that an improbable aspiration.

First, Carragher et al. (2015) point out that mental disorders are putatively distinct, yet they have a high level of comorbidity. The concept of comorbidity is often used in two ways: it can mean that a person meets the diagnostic criteria of more than one disorder, or that if someone can be classified as having a mental disorder, that person has more than one disorder with a higher probability than would be expected by chance alone (Lilienfeld, Waldman, & Israel, 1994). It is especially the latter meaning of the term that is both evident and problematic in the case of mental disorders. For example, in a nationally representative study of comorbidity of English-speaking adults in US, over 70 % of connections between 19 common mental and substance abuse disorders were positive and statistically significant (Kessler, Chiu, et al., 2005). On a more detailed level, some disorders co-occur more commonly than others, and in some cases comorbidity between specific disorders is rather the rule than an exception. As another example of this from the same nationally representative data from US, around two thirds of those with a DSM-IV diagnosed major depressive disorder also met the criteria of at least one another diagnosis when the depression had lasted at least 12 months (Kessler et al., 2003). The rates were the highest for anxiety disorders and then for impulse control and substance abuse disorders, with co-occurrence percentages being 57.5 %, 16.6 % and 8.5 %, respectively. This indicates that assumedly distinct categories are significantly overlapping and not, in fact, distinct.

Another related issue with categorical classifications lies with heterogeneity of symptoms within a single disorder, as pointed out by Carragher et al. (2015). The diagnostic categories of categorical mental disorders are polythetic in nature: they are diagnosed on the basis of various symptoms, but one does not need to have all the symptoms associated with a certain disorder in order to be diagnosed with the disorder. This brings about heterogeneity within disorder categories, because two people diagnosed with the same disorder might share only few common symptoms. To illustrate, in the case of DSM-5 major depressive disorder, two patients might share only one common symptom (American Psychiatric Association, 2015). Despite this, it is often thought that a single category describes a single mental disorder. This is a problematical assumption if several seemingly separate phenomena map into a single disorder category or if one phenomenon underlies several categories.

In the exemplary case of depression, there is evidence of both. In a study examining the factor structure of common measures developed for tools in diagnosing depression, it was found that the number of factors varied from three to six (Fried et al., 2016). This suggests that the symptoms of depression are multidimensional rather than unidimensional. On the other hand, research suggests that some factors, such as executive functions, underlie several mental disorders (Snyder, Miyake, & Hankin, 2015), including major depressive disorder (MDD) (Cotrena, Branco, Shansis, & Fonseca, 2016), yet executive functions are not explicitly included in the diagnostic criteria of MDD. This hints that current diagnostic categories might not include some relevant aspects integral to psychopathology.

Lack of validity brings about consequences both theoretical and practical in nature. If people having very different problems are lumped into one category when they should be split into several or vice versa, then it can be argued that the categorical classification in question is not able to claim the benefits by which the categorical classification systems are often defended: easing clinical work and aiding research (see Fried, 2015).

This criticism does not mean that current categorical classification systems do not have their merits. In fact, as Lilienfeld (2014) points out, the DSM-5 has made considerable efforts to take into account contemporary research. For example, it has deleted some problematic diagnostic categories, such as schizophrenia subtypes, which have shown poor validity, poor stability over time, poor interrater reliability and little utility (Reddy, Horan, & Green, 2014). Also, DSM-5 acknowledges the potential of some dimensional models by mentioning a hybrid dimensional-categorical model

for diagnosing personality disorders under “Emerging Measures and Models” – a section which is aimed at improving awareness and guiding research on potentially beneficial topics (APA, 2013).

Despite these efforts, many researchers have turned to dimensional approaches of mental disorders, which can avoid some of the issues categorical classifications face. For example, latent factors formed of an extensive number of measured symptoms and consisting of few, broad-band factors, can be understood as a way for taking account the pervasive comorbidity and correlation of different mental disorders (Eaton, South & Krueger, 2010). In a similar manner, dimensional models avoid problems of heterogeneity by not being polythetic.

1.2. The Proposed Latent Structures of Psychopathology

Numerous factor models underlying psychopathology have been proposed with varying methodologies and samples. Especially with regards to children and adolescent populations, a model consisting of two factors – internalizing and externalizing – has been used for decades to model childhood psychopathology (Achenbach, 1966; Achenbach & Edelbrock, 1978). This model has been often dubbed as the two-factor model, and it exhibits both good fit and stability over time (for review, see Eaton, Rodriguez-Seijas, Carragher, & Krueger, 2015), as well as invariance across genders, cultures, age and sexual orientation (Carragher et al., 2015). The factors have been interpreted as ways to explain comorbidity across psychiatric diagnoses or symptoms, as underlying core psychopathological processes (Caspi et al., 2014) and even as reflections of genetic etiology (Krueger, 1999).

The internalizing factor denotes liability to undergo depressive disorders and various anxiety disorders, such as generalized anxiety disorder, panic disorder, specific phobias, eating disorders and bipolar disorders, while the externalizing factor does the same for substance abuse disorders and disorders related to problems of behavioural, antisocial, or impulsive kind. Some disorders, such as borderline personality (Eaton et al., 2011; Forbush & Watson, 2013) and post-traumatic stress disorder (Carragher et al., 2015) seem to cross-load for both factors, which indicates heterogeneity in the disease categories. The internalizing factor is sometimes further divided into highly correlating fear and distress subfactors, but whether this bifurcated model exhibits better model fit compared to the two-factor model remains an open question (Eaton et al., 2015). Although the factors in the two-factor and the bifurcated models are deemed distinct, a correlation between the factors is often allowed. The correlations appear to be from moderate to strong: studies have

typically shown a moderate correlation of .4–.6 between the internalizing and externalizing factors (Forbush & Watson, 2013; Krueger, 1999; Patalay et al., 2015) and as demonstrated by Krueger (1999), a strong connection ($r = .73$) between the subfactors of internalizing.

Sometimes other similar factors are added to the models as well. The most common of these factors is the psychotic or the thought disorder factor. The principle remains the same as with the two-factor and bifurcated factor models: three correlating factors explain comorbidity across various symptoms and disorders. The psychotic factor is often omitted for practical reasons, since commonly research focuses on more common forms of psychopathology and most psychiatric surveys do not assess psychotic symptoms (Caspi et al., 2014). However, models containing the thought disorder factor have exhibited good fit (Kotov et al., 2011; Markon, 2010). Bipolar disorders and manic episodes seem to cross-load on the psychotic factor as well as the distress subfactor of internalizing, which is also theoretically plausible given the depressive–psychotic nature of bipolar disorders (for review, see Carragher et al., 2015). Similarly to the correlations of internalizing and externalizing factors, the thought disorder factor exhibits a significant correlation of around 0.6 with the internalizing factor (Wright et al., 2013).

1.2.1. The general factor of psychopathology.

Despite the good statistical fit of all the aforementioned models, the correlations among the factors could indicate the existence of a higher-level factor or several factors. This observation has generated research on a possible general psychopathology factor, which is sometimes called the p factor. A close analogy to the p factor is the general intelligence factor (g), which putatively explains why some people do well on a wide range of cognitive tasks. If understood via this analogy, the p factor could explain individual differences on a wide range of separate psychopathological symptoms.

The p factor has also been interpreted in more theoretical detail. For example, in a longitudinal study by Caspi et al. (2014) it was suggested that the p factor “summarizes individuals’ propensity to develop any and all forms of common psychopathology”. In the same study this interpretation seems to be backed by the association of general psychopathology and psychiatric history in family, as well as its association to high neuroticism, low conscientiousness and agreeableness and its ability to predict life impairment and compromised brain functionality. The p factor can also be interpreted as an indication of severity of psychopathology, for a rather obvious reason: since the p

factor is usually modelled with psychopathological symptom measures and symptom items generally load positively to the p factor, higher scores on the p factor reflect greater symptoms. This interpretation has been suggested by several researchers (cf. Arrindell et al., 2017; Caspi et al., 2014; Haltigan, 2019).

It should be noted that distinguishing a liability to psychopathology from the severity of psychopathology can be difficult, since phenomena such as psychopathological symptoms, criminal activity, other forms of life impairment and the use of substances can be plausibly interpreted as indications for both. However, the different interpretations might have their theoretical and possibly practical differences: for example, conceptualizing the p factor as a susceptibility factor could direct research into how negative life outcomes could be avoided by reducing the p factor, while the severity perspective could encourage clinicians into utilizing scales that measure the p factor in order to gain vital information about the depth of a patient's difficulties.

There is further evidence for all the theoretical interpretations mentioned above. First, the p factor explains comorbidity in itself by accounting for variance across different psychopathological disorders or symptoms. An early-emerging p factor is also able to prospectively predict school and global functioning, which indicates that the p factor acts as a susceptibility to negative life outcomes (Lahey et al., 2015). In support of the severity interpretation, De Raykeer et al. (2018) found that the p factor predicted suicide attempts in adults in a relatively short time period of three years. This indicates that higher levels of general psychopathology indeed reflect higher severity of psychopathology, not just susceptibility to develop mental disorders in distant future. Conway et al. (2019) found similar results with the severity of p factor predicting suicidality and self-injury during the past year.

In addition to having several possible interpretations, the p factor can also be formed in several models. For example, a model could consist only of the p factor. In that scenario, all items should load to the p factor. However, research suggests that a one-factor model consisting of the general psychopathology factor usually exhibits less than acceptable goodness-of-fit (Arrindell et al., 2017; Caspi et al., 2014; Patalay et al., 2015). A classic bifactor model consisting of both, the p factor and internalizing and externalizing (and perhaps thought disorder) factors, on the hand, often fits the data even better than various correlated-factors models without a general factor (Arrindell et al., 2017; Carragher et al., 2016; Lahey et al., 2012; Patalay et al., 2015). As such, all items should load to the p factor and either to the internalizing, externalizing or thought disorder factors, and by convention the model is usually fully orthogonal. If the bifactor model is formed this way, the role

of internalizing and externalizing factors can be understood as accounting for variability not fully accounted by the p factor (van Bork, Epskamp, Rhemtulla, Borsboom, & van der Maas, 2017). As demonstrated by Caspi et al. (2014), Lahey et al. (2012) and Patalay et al. (2015), bifactor models often show superior fit compared to correlated two- or three-factor models.

However, the classic bifactor model described above is not the only bifactor model that can be formed, and it is worth noticing that different bifactor models can have radically different interpretations. For example, the internalizing and externalizing factors can be allowed to correlate, or the p factor can be understood as a higher-order factor for the lower-level factors, such as internalizing, externalizing and thought disorders (see Blanco et al., 2015; van Bork et al., 2017). In the higher-order bifactor model the lower-level factors load onto the p factor while some of the observed variables, such as questionnaire items, load to the lower-level factors. In this case, the lower level factors explain the variance in their indicator items, while the p factor explains the variance that the lower-level factors share (van Bork et al., 2017).

As pointed out by van Bork et al. (2017), however, because model fit is measured by comparing observed variance–covariance matrix to the theoretical matrix postulated in the model, different higher-order and bifactor models are statistically highly similar and usually exhibit highly similar fit indexes. If fit indexes were the sole reason to prefer one model over others, this would result in a large risk of biased conclusions: small differences in the studied samples could lead one higher-order or bifactor model to emerge more fitting than the rest. This means that even a small sampling variability could lead to massively different theories of psychopathology, if models were compared based on just their fit indexes.

Focusing on the fit indexes is further complicated by bifactor models' tendency to fit virtually any data. Bonifay and Cai (2017) demonstrate that bifactor models have a proclivity to fit the data even when the data follows a random pattern. This indicates that while bifactor models are successful in terms of fit indexes, this could be a symptom of overfitting and ability to model noise (Bonifay, Lane, & Reise, 2017).

Partly because of these concerns, it is beneficial to examine different bifactor model properties above and beyond model fit, such as the external validity and practical applications of different models. This does not mean that goodness-of-fit of the models should not be used as evidence when comparing models – rather it means that a significant part of what makes one model better than others lies in its usefulness. Despite this, attempts to compare the usefulness of different models of psychopathology are still scarce. For example, if such models were able to predict how and what

kind of psychopathology at an early age affects later negative life outcomes, there might be a possibility to prevent such outcomes before they manifest. As a demonstration of this, it has been observed that externalizing disorders in preadolescence predict substance use in adolescence (King, Iacono, & McGue, 2004). Since early substance use can continue into adulthood (e.g. Jordan & Andersen, 2017) and lead into other negative consequences, interventions to underlying psychopathology could have highly impactful results. However, to the knowledge of the author of this study, similar research as in the study of King et al. (2004) has not been conducted with dimensional models of psychopathology.

1.3. Substance Use

On a general level, categorical diagnostic classes of substance use disorders (SUDs) are highly comorbid with other mental disorders (Degenhardt, Hall, & Lynskey, 2001; Kessler, Chiu, et al., 2005). This type of comorbidity can be called “heterotypic comorbidity”, since in it disorders from two different diagnostic groupings co-occur. There is also “homotypic comorbidity” in different substances, because often the users diagnosed with one substance-related disorder are more likely to exceed the criteria for other DSM-IV defined substance abuse or substance dependence disorders as well (Degenhardt et al., 2001). When it comes to diagnostic classes of substance use disorders, it should be noted that in DSM-5, unlike in DSM-IV, there is no distinction between substance abuse and substance dependence disorders but combines them into single category of SUDs. These DSM-5 defined SUDs vary in severity, ranging from mild, moderate to severe on the basis of the number of different substance use related symptoms.

The use of substances such as tobacco, alcohol and drugs cause considerable stress to affected individuals, their families and the entire society. Tobacco alone causes around 5 million annual deaths worldwide (Davis, Wakefield, Amos, & Gupta, 2007), whereas excessive alcohol consumption is one of the leading causes for premature mortality around the world (Leon et al., 2007; Rehm et al., 2007; Stahre, Roeber, Kanny, Brewer, & Zhang, 2014). Drug use, on the other hand, can cause several significant negative consequences such as unemployment, poor health and mental disorders (Das, Salam, Arshad, Finkelstein, & Bhutta, 2016).

In the development of SUDs or nicotine dependency, adolescence is a critical period. Nationally representative studies suggest that 50 % of SUDs begin by the age of 15–20 (Kessler, Berglund, et al., 2005; Merikangas et al., 2010). Early substance use can also continue into adulthood and further

develop into a SUD: research suggests that teens or children that use substances before the age of 14 have a higher risk of developing a substance dependence (Jordan & Andersen, 2017). There is also evidence that people who start using cannabis by the age of 17 are more likely to use, abuse or get dependent on drugs, and the same applies for alcohol dependence (Lynskey et al., 2003). It also has been estimated that each year early use of drugs is delayed, the risk of a lifelong drug abuse or dependency is decreased by around 4–5 % (Grant & Dawson, 1998).

From a dimensional perspective, the high homotypic comorbidity between different substance use disorders suggests that there could be one or several factors that help to explain the comorbidity. Indeed, studies have frequently found that there might be an underlying common factor behind SUDs (cf. Hasin, Fenton, Beseler, Park, & Wall, 2012; Huba, Wingard, & Bentler, 1981). However, the latent factor could also encompass more than just SUDs. It has also been noted that substance use often co-occurs with tendencies for impulsivity and aggression as well as antisocial personality, which has led some researchers to suggest that these tendencies form a latent factor of externalizing (Krueger, Markon, Patrick, Benning, & Kramer, 2007). This is supported by confirmatory factor analyses that have found that SUDs often load highly to externalizing as well as the p factor (Greene & Eaton, 2017; Ignatyev, Baggio, & Mundt, 2019).

Due to the critical nature of preadolescence in development of SUDs, the burden of SUDs and the high comorbidity of SUDs and other mental disorders, research on predicting substance use and abuse before it even starts could yield important real-life applications. These reasons also make the topic an ideal candidate for examining the usefulness of dimensional models of psychopathology.

1.4. Research Questions

The aim of this study is to study whether several factor models of psychopathology that have been proposed in previous literature are viable in terms of statistical fit, and which models emerge useful in predicting early experimentations of alcohol, tobacco, marijuana, glue and other drugs. More specifically, the models examined are the oblique two-factor model, the unidimensional general psychopathology factor model and the orthogonal bifactor model.

2. Methods

2.1. Participants

The data of this study is from the UK Household Longitudinal Survey, a longitudinal study which consists of members of approximately 40 000 households in United Kingdom. The data collection started in 2009 and has taken and will take place in 8 waves with the last wave being collected in 2019. The participants of this study participated in the initial measurements in waves 1 and 3 (collected in 2009–2010 and 2011–2012, respectively) and in the follow-up measurements in waves 4 and 6 (in 2012–2013 and 2014–2015, respectively).

The target sample of this study consisted of 3648 (49.5 % female) preadolescents aged from 10 to 12 years (mean = 11.0, sd = 0.77) at the time of the initial measurement. Due to a drop-out rate, at second phase of the study the sample consisted of 2244 participants aged from 12 to 16 years (mean = 13.91, sd = 0.91).

2.2. Measures

The psychiatric symptoms were measured by the Strengths and Difficulties Questionnaire (SDQ), which is a widely used method for assessing the behaviour, emotions and relationships of 4–16-year-olds. SDQ can be used for clinical screening or assessment as well as for research (Goodman, 2001). SDQ boasts good psychometric properties (Goodman, Meltzer, & Bailey, 1998), and it has been successfully used to model latent factors of psychopathology (Carragher et al., 2016; Patalay et al., 2015).

In this study, 15 items from the SDQ were used, although the original questionnaire consists of 25 items in total. Originally the 25 items were intended to form 5 subscales (emotional problems, peer problems, conduct problems, hyperactivity and prosociality), each consisting of 5 items. Example items include questions such as “I worry a lot” and “I am often accused of lying or cheating”. In this study, the items forming the prosociality subscale were omitted, since they are not measuring psychiatric symptoms. The remaining four subscales also included 5 items in total which are reverse-worded and measure strengths instead of psychiatric symptoms. Because the correlation among these reverse-coded items and the other items assessing problems has been shown to be weak, the coefficient alpha and thus potentially cause problems to the factor solutions (Van De Looij-Jansen, Goedhart, De Wilde, & Treffers, 2011), these items were also omitted from the

analyses of this study. All SDQ are answered on a 3-point scale (1 = “Not true”, 2 = “Somewhat true”, 3 = “Certainly true”).

Of the 15 items used, internalizing was assessed with 8 items ($\alpha = .88$) and externalizing with 7 items ($\alpha = .85$). General psychopathology was assessed with all 15 items ($\alpha = .92$). The decision on which items were categorized as internalizing and which as externalizing was based on previous studies employing the same measure in study of factors underlying psychopathology (Carragher et al., 2016; Patalay et al., 2015).

The early use of alcohol, tobacco, cannabis, glue or solvent sniffing and any other illegal drugs was assessed with questions such as “Have you ever had an alcoholic drink? That is a whole drink, not just a sip.”, and in all items the participants had to choose from two choices: “Yes” and “No”.

2.3. Procedure

The participants took part at either waves 1 and 4 or 3 and 6, and the data from all waves was pooled. At the initial measurements at waves 1 or 3, the participants completed an extensive questionnaire concerning their life and answered the Strengths and Difficulties Questionnaire, among other items.

On the follow-up phase (waves 4 or 6) the procedure was identical except for the contents of the questionnaire – this time it included, among other items, questions about substance use. If and only if a person was given the questionnaire containing SDQ items at wave 1, then questionnaire containing the substance use items was administrated around 3 years later in wave 4; and similarly, if and only if the questionnaire containing SDQ was administrated at wave 3, substance use items followed around three years later in wave 6. At all waves the contents of the questionnaires also included items not relevant to this study.

2.4. Statistical Analyses

The analyses can be summarized into two steps. First, confirmatory factor analyses (CFAs) were conducted. In this step, the goodness-of-fit of the models and other model statistics – namely omega, omega hierarchical and H index – were also examined. In the second step, structural

equation models were conducted to predict whether adolescents had had early experiences with substances. Both the SEMs and the CFAs were conducted using the R package lavaan version 0.6-5. SDQ items and substance use items were treated as categorical variables, given the skewness of the variables and that the former has only three and the latter only two levels. The estimation method for both the CFAs and SEMs was diagonally weighted least squares, because it fares better than maximum likelihood equations when data is ordinal (Li, 2016).

2.4.1. Confirmatory factor analyses, model fit and properties of the models.

Underlying structure of psychopathology was first modelled with three confirmatory factor analyses. The models were the two-factor model consisting of correlating internalizing and externalizing factors; the one-factor model consisting of the p factor model; and the orthogonal bifactor model consisting of all three: the internalizing and externalizing factors and the p factor. The models are presented visually in Figure 1, Figure 2 and Figure 3. After that, the goodness-of-fit indexes of the models were examined. While there is no clear consensus regarding strict cut-offs in fit indexes, often RMSEA values below .05 are deemed reasonably good in terms of fit, whereas values below .1 should be rejected (Brown, 2006). Similarly, comparative fit index (CFI) and Tucker–Lewis index (TLI) values above .95 can be understood to signify good fit and values between .9 and .95 can be called “acceptable”, whereas values below .9 should be rejected (Brown, 2006). In the case of SRMR, it has been suggested that .08 is required for a relatively good fit (Hu & Bentler, 1999). For the purposes of this study, RMSEA and SRMR values below .1 and .08, respectively, and CFI and TLI values above .9 are considered acceptable.

Since bifactor models have a tendency to overfit data (Bonifay et al., 2017), it is important to evaluate bifactor models with methods that have been specifically designed for evaluating bifactor models. Thus, properties of the CFA models in this study, especially the bifactor model, were also examined with omega, omega subscale (omegaS), omega hierarchical (omegaH), omega hierarchical subscale (omegaHS), explained common variance (ECV), percentage of uncontaminated correlations (PUC) and H indices. Omega subscale is a tool for estimating the internal reliability of a factor analytic model, and it can be applied to both: the whole model or the subscales of the model, hence the distinction to omega and omega subscales (Rodriguez, Reise, & Haviland, 2016a). Omega resembles the standard coefficient alpha that is widely used in psychological research. However, there are two differences: omega is based on factor loadings

rather than observed variances and covariances, and omega is more suitable in situations where loadings vary (Rodriguez, Reise, & Haviland, 2016b).

Along with omega, omegaH and omegaHS are informative indexes in evaluating bifactor models. OmegaH is used to estimate degree of systematic variance in raw total scores that is attributable to individual differences in the p factor, whereas omegaHS refers to the proportion of systemic variance that is unique to group factors (Rodriguez et al., 2016a). When omegaH is high (e.g. a cut-off $> .80$ can be used), such a significant proportion of reliable variance is attributable by the general factor that the scores can be viewed as essentially unidimensional (Rodriguez et al., 2016a).

ECV is another tool that can be used to estimate the degree of unidimensionality. When the general factor explains a large proportion of variance, value of ECV is high, which indicate a strong general factor (Reise, 2012). In extreme cases, when for example $ECV > .70$ or $> .80$, the data can be declared as essentially unidimensional. However, ECV values are moderated by PUC, which is an indicator for bias that would result in trying to fit an unidimensional model when the data would be better suited for a multidimensional model (Reise, Scheines, Widaman, & Haviland, 2013). Simply put, PUC in conjunction with ECV can be used as information about whether unidimensional model is appropriate, or whether doing so would lead to a large bias. H coefficient, on the other hand, is a measure of construct replicability or reliability, which measures the correlation between a factor and an optimally weighted item composite (Rodriguez et al., 2016b). When the value of H coefficient is high (e.g. $> .80$), the score indicates stability across studies, whereas a low value indicates that a factor is not well defined and might not be replicable (Rodriguez et al., 2016a). Rodriguez et al. (2016a) also suggest that when the H value of a factor is less than .70, the factor is not worth specifying. In this study, the omegaH, omegaHS, ECV and PUC were calculated only for the bifactor model, since they were used as tools in evaluating the appropriateness of the bifactor model. The examination of these indices was conducted by using Excel-based calculators provided by Dueber (2017) and Hammer (2016).

2.4.2. Structural equation models.

After the examination of the properties of the factors, whether adolescents had had early experiences with various substances was predicted. The predictions were conducted with multiple probit regressions in structural equation models (SEMs) and all regression analyses included sex, age in phase 4 or 6, ethnicity and family income as covariates. Probit regression is an alternative to logistic regression, which is not supported by lavaan version 0.6-5. The regression approach was a

one-step “forced approach”. The SEMs utilized the same latent factor structures as in those CFA models that had proven to be at least acceptable in terms of statistical fit. Those models that did not reach acceptable levels of fit, were not further used in SEMs.

There were several reasons for conducting both, separate CFAs and SEMs, instead of either just conducting SEMs or CFAs and factor score regression. The reasons were three-fold: first, by conducting separate CFAs, the properties of the latent factors could be examined more accurately without regressions, which could affect the fit indexes. Secondly, the SEMs were executed with a significantly smaller sample size ($n = 1610$) than the CFAs ($n = 3410$) due to the drop-out rate between the phases, so by modelling the latent structures without the regressors the risk of bias was lower. Thirdly, another source of bias was minimized when conducting regressions as a part of the SEMs instead of factor score regression. This is because there is an infinite number of ways to score individuals from factor loadings (Grice, 2001) and because the common methods for estimating factor scores and conducting factor score regression based on them produce more biased regression coefficients than regression analyses calculated as a part of SEM produce (Devlieger, Mayer, & Rosseel, 2016).

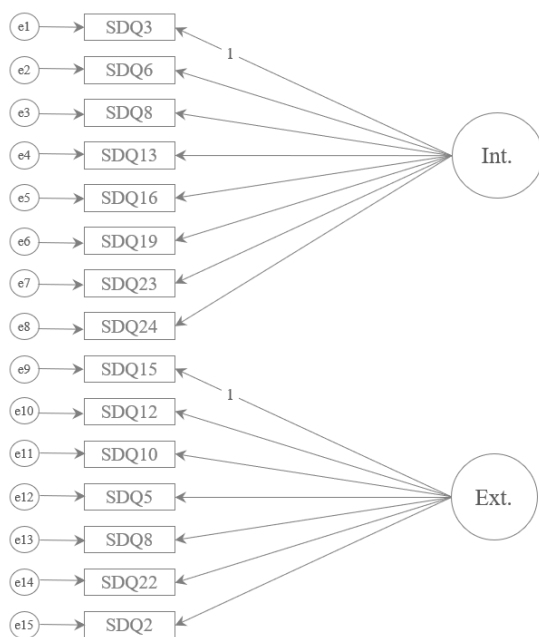


Figure 1. The two-factor model. Int. = Internalizing, Ext. = Externalizing.

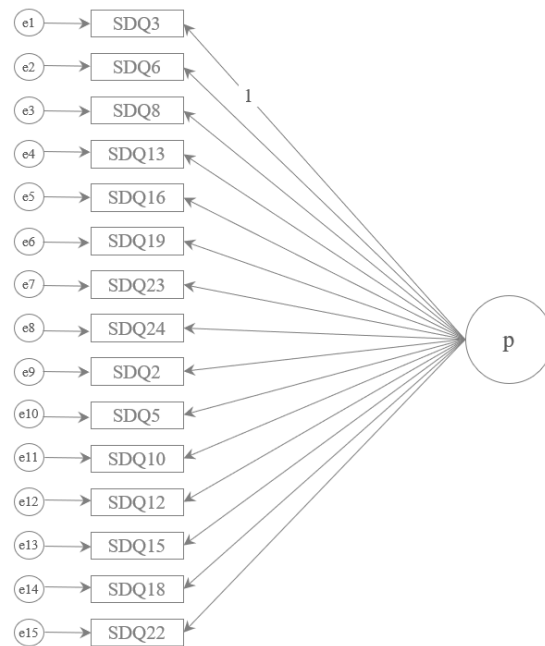


Figure 2. The p factor model. p = The p factor (the general psychopathology factor).

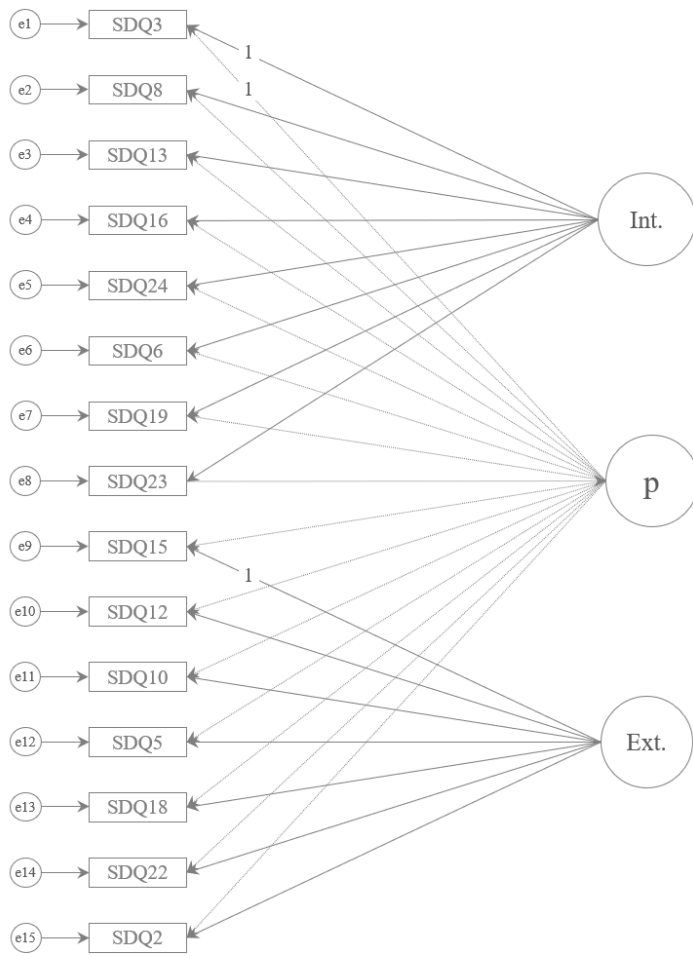


Figure 3. The bifactor model. p = The p factor (the general psychopathology factor), Int. = Internalizing, Ext. = Externalizing.

3. Results

3.1. Confirmatory Factor Analyses and Model Statistics

Based on the goodness-of-fit indexes, the two-factor model fit the data well: $\chi^2(89, N = 3437) = 956.56, p < .001$, CFI = .959, TLI = .951, RMSEA = .053, 90 % confidence interval = [.050, .056], SRMR = .059. There was moderate correlation between the latent factors ($r = .630$). The bifactor model also fit the data well and even better than the two-factor model: $\chi^2(75, N = 3437) = 367.441, p < .001$, CFI = .986, TLI = .980, RMSEA = .034, 90 % confidence interval = [.030, .037], SRMR = .041. A likelihood ratio test was calculated to compare the fit of the two-factor and the bifactor models. The results showed that the difference was statistically significant, $X^2(df = 14) = 589.12, p < .001$. The unifactorial model consisting of a sole p factor, however, did not fit the data acceptably:

$\chi^2(90, N = 3437) = 1911.799, p < .001, CFI = .913, TLI = .899, RMSEA = .077, 90 \% \text{ confidence interval} = [.074, .080], SRMR = .084$. This means that based on TLI and SRMR values, the one-factor model exhibited less than acceptable fit and was not used in further SEMs to predict future substance use.

Standardized factor loadings, their means and standard deviations as well as the values of omega, omega hierarchical and H index are presented in Table 1. For the two-factor and one-factor models, all of the standardized factor loadings were positive and above 0.3, with only one loading being below 0.4. For the bifactor model, however, especially on the externalizing factor the standardized factor loadings were lower, with 2 out of 7 being non-significant and a total of 5 being below 0.2. Due to convergence issues, the factor loading of the SDQ item number 15 to the externalizing factor was set to 1.0 in the CFAs that had an externalizing factor, even though usually by convention the first item in numerical order is fixed to 1.0 if one of the factor loadings to a latent factor is fixed in the first place.

The inspection of omegaH in the bifactor model also revealed that most of the variance of raw total scores was attributable to the general factor, while only a small portion was attributable to the group factors after taking into account the variability attributed to the general factor (cf. Reise, Bonifay, & Haviland, 2013). To be more specific, the percentage of reliable variance caused by internalizing and externalizing in total scores can be calculated by simply by subtracting the omegaH score (.715) of the p factor from its omega score (.881) (Rodriguez et al., 2016b). This results in 16.6 % of reliable variance being attributable to internalizing and externalizing subscores. In a similar manner, Rodriguez et al. (2016b) instruct that the proportion of reliable variance that can be attributed to the general factor can be calculated by dividing the p factor's omegaH score (variance due to the p factor) with its omega score (variance due to all factors in the model). This results in almost all reliable variance (81.1 %) being attributable to the general factor. These results are further supported by the results concerning the ECV in general factor (ECV = .63) and PUC (PUC = .53) in the bifactor model. Reise et al. (2013) suggest that when PUC is lower than .80, ECV of a general factor is greater than .60 and omegaH is greater than .70, the bifactor model can be interpreted as unidimensional. In this study, these criteria were met.

Table 1.

Standardized factor loadings and indices of omega, omega hierarchical and H for the two-factor model, the one-factor model and the bifactor model.

*Factor loadings are statistically significant with $p > .001$, except for those loadings with special markings. * = $.05 > p > .01$, x = not statistically significant*

<u>SDQ item</u>	The two-factor model		The one-factor model	The bifactor model		
	<u>Internalizing</u>	<u>Externalizing</u>	<u>p factor</u>	<u>p factor</u>	<u>Internalizing</u>	<u>Externalizing</u>
3. I get a lot of headaches, stomach-aches or sickness	.45		.41	.35	.26	
8. I worry a lot	.61		.53	.33	.61	
13. I am often unhappy, down-hearted or tearful	.78		.69	.55	.51	
16. I am nervous in new situations. I easily lose confidence	.57		.51	.39	.41	
24. I have many fears, I am easily scared	.55		.48	.31	.52	
6. I am usually on my own. I generally play alone or keep to myself	.49		.44	.35	.31	
19. Other children or young people pick on me or bully me	.63		.56	.44	.43	
23. I get on better with adults than with people my own age	.43		.39	.30	.30	
15. I am easily distracted, I find it difficult to concentrate		.70	.65	.67		.18
12. I fight a lot. I can make other people do what I want		.59	.54	.62		-0.05x
1. I am constantly fidgeting or squirming		.69	.64	.57		.67
5. I get very angry and often lose my temper		.69	.64	.72		-0.03x
18. I am often accused of lying or cheating		.64	.60	.68		-0.10*
22. I take things that are not mine from home, school or elsewhere		.53	.50	.56		0.10*
2. I am restless, I cannot stay still for long		.60	.54	.46		.53
Mean	.57	.63	.54	.49	.42	.13
Standard deviation	.11	.07	.09	.15	.13	.30
Omega/omegaS	.79	.83	.80	.88	.79	.85
OmegaH/omegaHS				.72	.44	.05
Explained common variance (ECV)				.63	.25	.13
H index	.82	.83	.87	.86	.67	.56

3.2. Structural Equation Models and Prediction of Substance Use

Two SEMs were conducted to predict substance use, namely has the person ever tried alcohol, tobacco, glue or solvents, cannabis or other drugs. By the time the regressions were conducted, 8.0 % had already tried smoking, 33.1 % had tried alcohol, 1.7 % had sniffed solvent or glue, 3.1 % had tried marijuana and finally, 0.5 % had tried other drugs. The latent structures used for predictions were the two factors in the two-factor model and the three factors in the bifactor model. Similar to the CFAs, the factor loading of the SDQ item number 15 to the externalizing factor was set to 1.0 in both models due to convergence issues.

Adding the regressions reduced the goodness-of-fit of both models. In the case of the two-factor SEM model, CFI and TLI seemingly decreased, while RMSEA and SRMR increased: $\chi^2(214, N = 1610) = 959.59, p < .001$, CFI = .930, TLI = .938, RMSEA = .047, 90 % confidence interval = [.044, .050], SRMR = .069. The bifactor model $\chi^2(195, N = 1610) = 652.56, p < .001$, CFI = .957, TLI = .958, RMSEA = .038, 90 % confidence interval = [.035, .041], SRMR = .057. Similarly, the goodness-of-fit also changed in the bifactor model by adding the regressions, with CFI and TLI decreasing and RMSEA and SRMR increasing: $\chi^2(195, N = 1610) = 652.56, p < .001$, CFI = .957, TLI = .958, RMSEA = .038, 90 % confidence interval = [.035, .041], SRMR = .057. This means that the fit of both models worsened slightly by adding the multiple regressions but remained acceptable in both models.

The regression coefficients and other regression test statistics for multiple probit regressions are presented in Table 2. In the two-factor model, the externalizing factor was a statistically significant and positive predictor of the use of all types of substances, as was the p factor in the bifactor model.

Table 2.

Multiple probit regression results for the two-factor model and the bifactor model.

		Unstand.	Stand.	z	p
<u>The two-factor model</u>					
Smoking	Internalizing	-0.27	-0.11	-1.81	.07
	Externalizing	0.55	0.37	5.86	< .001
Alcohol	Internalizing	-0.35	-0.14	-3.10	.002
	Externalizing	0.33	0.20	4.68	< .001
Glue / solvent	Internalizing	-0.23	-0.09	-0.97	.33
	Externalizing	0.43	0.28	2.92	.004
Cannabis	Internalizing	-0.38	-0.15	-1.85	.06
	Externalizing	0.58	0.36	4.55	< .001
Other drugs	Internalizing	0.08	0.03	0.61	.54
	Externalizing	0.51	0.33	6.75	< .001
<u>The bifactor model</u>					
Smoking	p factor	0.95	0.30	7.21	< .001
	Internalizing	-0.44	-0.10	-1.89	.06
	Externalizing	0.09	0.02	0.27	.79
Alcohol	p factor	0.41	0.12	4.42	< .001
	Internalizing	-0.52	-0.12	-2.84	.004
	Externalizing	0.08	0.01	0.31	.76
Glue / solvent	p factor	0.74	0.23	3.90	< .001
	Internalizing	-0.40	-0.09	-1.13	.26
	Externalizing	-0.02	0.00	-0.03	.98
Cannabis	p factor	0.93	0.28	5.52	< .001
	Internalizing	-0.69	-0.15	-2.11	.04
	Externalizing	0.18	0.03	0.41	.68
Other drugs	p factor	1.38	0.43	10.81	< .001
	Internalizing	-0.31	-0.07	-1.42	.16
	Externalizing	-0.78	-0.14	-2.29	.02

Note. All models included the age, sex and race of the participants and family income as covariates. Unstand. = unstandardized regression coefficients, Stand. = standardized regression coefficients.

4. Discussion

This study examined the structure of psychopathology in preadolescence and the viability of competing latent factor models in predicting substance use in adolescence. Both the two-factor model and the bifactor model, in which all items loaded to general psychopathology, were viable solutions for the underlying structure of psychopathology in terms of fit, whereas the unidimensional general factor model was not. Both viable models were also able to predict future substance use.

In the two-factor model, more severe externalizing symptoms predicted all forms of early substance use, namely the use of cigarettes, alcohol, cannabis and any other illegal drugs as well as sniffing of glue or solvent, whereas in the bifactor model more severe general psychopathology managed the same. These results corroborate the findings that higher levels of externalizing are indeed linked to increased substance use (Greene & Eaton, 2017; Ignatyev et al., 2019; Krueger et al., 2007). It is notable that of all substances, experimentation with alcohol was the weakest prediction for both: the general factor in the bifactor model and the externalizing factor in the two-factor model. This is likely because at the time of the measurement around a third had already tried alcohol, and thus many adolescents who do not have particularly severe psychopathological symptoms had probably also experimented with it. On the other hand, more severe general psychopathology predicted the use of other drugs more strongly than the use of other substances, whereas in the two-factor model, more severe externalizing symptoms predicted the use of cigarettes more strongly than the use of other substances. This suggests that the use of highly hazardous drugs such as heroin, cocaine and methamphetamine reflects the severity of an individual's psychopathological symptoms more so than the use of other substances, such as alcohol, cigarettes or marijuana.

The strength of the predictions between the models could not be compared due an inherent difficulty in comparing the results of different probit or logistic regression models (Karlson, Holm, & Breen, 2012), but incipient research has suggested that correlated factors models are in many cases comparably good in predicting negative life outcomes, though there may be specific areas where one model performs better than others. Laceulle, Chung, Vollebergh and Ormel (2019) examined how different models of psychopathology predicted a variety of life outcomes, ranging from e.g. education level to use of cigarettes, alcohol or cannabis. The models that were analyzed in depth were a three-factor model consisting of correlating internalizing, externalizing and thought disorders and a bifactor model that consisted of a general psychopathology factor to which all items loaded in addition to the three group factors. The researchers found that higher levels of

externalizing in the correlated-factors model predicted the use of cigarettes, alcohol and cannabis better than the general factor in the bifactor model did, although generally neither of the models came out superior. Caspi et al. (2014) also studied the associations of two models underlying psychopathology with different types of life impairment, namely suicide attempt, psychiatric hospitalization, duration of social-welfare benefit use and convictions of violent crimes. They found that both examined models – which were structurally the same as the two in the study of Laceulle et al. (2019) – had factors that were substantially positively associated to all measured types of life impairment. While more severe general psychopathology was more strongly associated to most types of life impairment than any of the factors in the correlated factors model, the differences were small, and in the case of convictions of violent crimes, the association to externalizing in the three-factor model was slightly stronger than it was to the p factor in the bifactor model. In sum, these results suggest that although in general the models fare comparably well, in prediction of substance use the two-factor model might be more useful. However, more research is needed to verify or falsify this conclusion.

It is also worth noticing that while the bifactor model in this study did fit the data better than the two-factor model, bifactor models have a tendency to fit due to their greater ability to capture noise and random patterns (Bonifay & Cai, 2017; Bonifay et al., 2017). Therefore, the better goodness-of-fit of the bifactor model does not directly indicate that it reflects the structure of psychopathology better than the two-factor model. Nevertheless, especially in the case of the bifactor model, the good fit of the model is noteworthy due to the young age of the participants (mean = 11.02 years). This is younger than, to the knowledge of the author of this article, the average age in all but three studies that have found the p factor (Lahey et al., 2015; Martel et al., 2016; Olino et al., 2018), with one article coming close (Gomez et al., 2019). This corroborates the yet-scarce literature that modeling preadolescence psychopathology with a bifactor model can be viable. The lackluster fit of the one-factor model has also been established before (Arrindell et al., 2017; Carragher et al., 2016; Caspi et al., 2014; Martel et al., 2016; Patalay et al., 2015), which suggests that as with adults and adolescents, it is not feasible to model psychopathology in preadolescence with only an unidimensional structure.

Despite the unidimensional model not being viable, curiously enough, according to a heuristic proposed by Reise et al. (2013), the bifactor model in this study can be considered unidimensional due to the strength of the general psychopathology factor. In other words, although in principle the theoretical role of internalizing and externalizing group factors in the bifactor model is to help to explain aspects of psychopathology not comprehensively explained by the p factor (Bonifay et al.,

2017; van Bork et al., 2017), in this study their role in explanation was rather negligible. Especially the externalizing factor explained very little of the common variance or the systematic reliable variance that is independent of the general factor. In a similar manner, the group factors in the bifactor model should not be interpreted independently of the general factor (cf. Rodriguez et al., 2016a). This puts the theorist who is trying to choose the most theoretically sound model in a difficult spot: the unidimensional model is not feasible, yet the feasible bifactor model is primarily unidimensional – but in it the feasibility comes with the cost of including “methodological nuisance” factors, which cannot be given meaningful theoretical interpretations (cf. Bonifay et al., 2017). The two-factor model, on the other hand, can be more easily interpreted as consisting of internalizing and externalizing, since in it both factors were well-formed.

It should be kept in mind, however, that the results concerning the strength of the p factor have been mixed. In a study by Gomez et al. (2019) the findings did not support the unidimensionality of the examined bifactor models and, with the exception of internalizing in adolescence, all factors were well-defined and could be given meaningful interpretations. A study by Arrindell et al. (2017), however, did suggest very low levels of reliable variance attributable solely to the group factors in two separate bifactor models. According to another heuristic ($PUC > .80$) put forward by Reise et al. (2013), both bifactor models in the study could also be interpreted as unidimensional. A third study on the matter also concluded that a bifactor model examined in the study had interpretative issues similar to those in this study, since the p factor explained most of the variance, which led to unreliability of the group factors (Conway, Mansolf, & Reise, 2019). The mixed results go to show that bifactor derived indices should be calculated by default in research on bifactor models. This would aid in answering, for example, are the results replicable or specific to the used measures.

In addition to issues with interpretation, it is also difficult to apply the group factors to practice due to the low reliability and proportion of explained variance of the group factors, alongside with the fact that the group factors are not well defined. Researchers of Wechsler’s cognitive skills tests have warned that the subscales of Wechsler’s intelligence tests should be interpreted with caution (Kush & Canivez, 2019; McGill, Dombrowski, & Canivez, 2018) and a similar suggestion is in place with the group factors of the bifactor model, since the results in this study are similar to that of the Wechsler literature. This goes against the conclusions of Carragher et al. (2016), who examined structurally a very similar bifactor model partly with a same measure (SDQ) as in this study and suggested that in clinical practice both the p factor and the group factors should be assessed to “provide detailed information about a patient’s [risk] profile”. Carragher et al. (2016),

however, did not examine any of the bifactor indices utilized in this study, which could have affected the authors' conclusions.

The residualized nature of the group factors in the bifactor model is also a likely explanation to some counterintuitive findings in this study. Some items in externalizing in the bifactor model had negative or statistically non-significant factor loadings for externalizing and *lower* levels of externalizing predicted early experiments with other drugs. This could be a result of the relative strength of the general psychopathology factor: since it explains most of the common variance, the externalizing factor loses its original meaning. Conway et al. (2019), who had similar interpretative issues, arrived at similar conclusions concerning the group factors in their study. It should be noted, however, that lower levels of internalizing predicted early experiments with alcohol in both the bifactor and the two-factor model and early experiments with marijuana in the bifactor model. The association between marijuana and internalizing in the two-factor model was also very close to being statistically significant ($p = .06$). This suggests that those preadolescents higher in internalizing symptoms might be at a reduced risk of using substances, at least when it comes to alcohol and perhaps to cannabis as well, even though the severity of their psychopathological symptoms at large would make them more susceptible to use alcohol and cannabis at an early age in adolescence. While the internalizing factor in the bifactor model was not well-defined, the results from the two-factor model corroborate that the results with the bifactor model could be meaningful. It is also intuitively plausible: the internalizing items in SDQ reflecting introversion and problems with peers, which could limit adolescents' access to alcohol, as well as other substances.

4.1. Limitations

Since the SDQ does not assess psychotic symptoms, a model consisting of a thought disorder factor could not be examined. Including psychotic symptoms into a model of latent psychopathology would be important, however, due to the high comorbidity of psychotic symptoms and their economic consequences (Carragher et al., 2016; Caspi et al., 2014). It is also worth noticing that the participants' answers to both, the SDQ and the questions about substance use were self-reported. This could have biased the results by, for example, some participants not answering truthfully about their use of substances or about their difficulties measured by the SDQ. There is also a risk of bias due to the drop-out rate between the first and the second time of measurement.

On a different SDQ-related matter, due to the categorical nature of the SDQ items and items measuring substance use, direct comparison of models was difficult. As in logistic regression, in

probit regressions the proportion of explained variance (R^2) cannot be calculated. Some alternatives to the standard R^2 have been developed for logistic and probit regressions, but in general they are estimates of goodness-of-fit and do not purport to explain shared variance (Veall & Zimmermann, 1994). Also, lavaan version 0.6-5 does not support the calculation any type of pseudo- R^2 in the first place, which is why no pseudo- R^2 s were calculated in this study. Similarly, because the predictions were conducted with multiple probit regressions, reporting intuitively understandable effect sizes was not possible and the regression coefficients between the two-factor model and the bifactor model could not be compared straightforwardly by examining the strength of the standardized regression coefficients (Karlson et al., 2012). Karlson et al. (2012) explain that this is because in probit and logit models, adding or removing any variable affects the error variance of the whole model. This in turn will cause the regression coefficients to be on different scales, which makes direct comparison between different models unviable without further analyses. A method developed by Karlson et al. (2012) makes it possible to compare probit regression coefficients, but it could not be conducted in lavaan 0.6-5. On the other hand, goodness-of-fit indexes were calculated for SEMs that included the regressions, but in order to compare the regression coefficients or variance explained in competing models, future research might be better off opting for data that can be analysed with linear regression.

4.2. Conclusion

While the group factors in the bifactor model were not well-formed and could not be interpreted, the bifactor model seems like a possible way to model preadolescent psychopathology. The value of bifactor models of psychopathology often seems to boil down to the predictive power of the general factor, and this study indeed corroborated the general factor's ability to predict important outcomes, namely the use of different substances in adolescence. However, the two-factor model without the general factor was also successful in predicting substance use and was less problematic in terms of interpretation. Earlier research also suggests that externalizing in a correlated factors model, where a general factor does not take away systematic variance explained by externalizing, might be superior in predicting the use of substances (Laceulle et al., 2019). However, further research is needed to examine this. In sum, while the bifactor model has its merits in predicting life outcomes, the theoretically simpler correlated-factor alternatives seem to be at least comparably useful with regards to the use of substances. Future research should compare the practical utilities of different models of psychopathology and examine bifactor model indices, such as omega hierarchical and

explained common variance by default, so we can reach a more nuanced understanding of psychopathology.

5. References

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